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IN THE CLAIMS

1. (currently amended) A method of screening for the discovery of disease associated molecular targets for diagnostic or therapeutic intervention, the method comprising;

performing *in vivo* imaging <u>by dynamic contrast MRI</u> of diseased tissue <u>from an individual</u> to provide one or more *in vivo* images;

evaluating said in vivo image for imaging features;

obtaining a cellular sample from said diseased tissue <u>from said individual</u>, which sample corresponds to said imaging feature;

determining the expression of genes or gene products in said cellular sample;

comparing said expression in said cellular sample with a control tissue;

wherein genes or gene products upregulated in said cellular sample represent molecular targets for therapeutic or diagnostic intervention.

2. -3 (canceled)

- 4. (original) The method according to Claim 1, wherein said step of determining expression of genes comprises hybridization analysis of probes derived from mRNA present in said cellular sample.
- 5. (original) The method according to Claim 1, wherein said step of determining expression of gene products comprises proteomic analysis.
- 6. (original) The method according to Claim 1, wherein said control sample comprises cells from said diseased tissue, but spatially or temporally separated from said cellular sample.
- 7. (currently amended) The method according to Claim 1 A method of screening for the discovery of human disease associated molecular targets for diagnostic or therapeutic intervention, the method comprising;

performing *in vivo* imaging wherein said *in vivo* imaging is selected from the group consisting of MRI, MRS, nuclear scintigraphy, PET, CT, ultrasonography, optical imaging, infrared imaging, and x-ray radiography of diseased tissue from a human patient to provide one or more *in vivo* images; evaluating said *in vivo* image for imaging features;

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obtaining a cellular sample from said diseased tissue from said human patient, which sample corresponds to said imaging feature;

determining the expression of genes or gene products in said cellular sample;

comparing said expression in said cellular sample with a control tissue;

wherein genes or gene products upregulated in said cellular sample represent molecular targets for therapeutic or diagnostic intervention.

8-22 (canceled)

23. (new) The method according to Claim 1, wherein said individual is ahuman.